STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source:

Date Processed by STIC:

10/542,239

07/25/2005

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
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Revised 01/24/05

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SUGGESTED CORRECTION ASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY Wrapped Nucleics The number/text at the end of each line "wrapped" down to the next line. This may occur if your file Wrapped Aminos - was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will _____prevent "wrapping." Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces. The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers: Misaligned Amino Numbering use space characters, instead. The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please Non-ASCII ensure your subsequent submission is saved in ASCII text. contain n's or Xaa's representing more than one residue. Per Sequence Rules, Variable Length each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing. A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid PatentIn 2.0 __. Normally, Patentin would automatically generate this section from the "bug' previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences. missing. If intentional, please insert the following lines for each skipped sequence: Skipped Sequences Sequence(s) (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (OLD RULES) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION: SEQ ID NO: X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences. Skipped Sequences Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence. (NEW RULES) <210> sequence id number <400> sequence id number 000 Use of n's and/or Xaa's have been detected in the Sequence Listing. Use of n's or Xaa's Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. (NEW RULES) In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents. Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or Invalid <213> scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or Response is Artificial Sequence missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> Sequence(s) Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220 to <223 section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules) Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file. Patentin 2.0

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resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence

"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid

listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

"bug"

Misuse of n/Xaa



PCT

RAW SEQUENCE LISTING DATE: 07/25/2005
PATENT APPLICATION: US/10/542,239 TIME: 11:54:35

Input Set : A:\1181-8 PCT US.SeqList.txt
Output Set: N:\CRF4\07252005\J542239.raw

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              Bradu, Stefan M.
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              Hannan, Raquib
              Pincus, Matthew R.
      8 <120> TITLE OF INVENTION: PANCREATIC CANCER ASSOCIATED ANTIGEN, ANTIBODY THERETO, AND
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     11 <130> FILE REFERENCE: 1181-8 PCT US
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C--> 14 <141> CURRENT FILING DATE: 2005-07-15
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     16 <150> PRIOR APPLICATION NUMBER: PCT/US2004/001196
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     17 <151> PRIOR FILING DATE: 2004-01-16
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57 Invalid Response
See Hem # 11 on
See From Summery Sheet-
Error Summery Sheet-
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     20 <151> PRIOR FILING DATE: 2003-01-17
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DATE: 07/25/2005 TIME: 11:54:35

PATENT APPLICATION: US/10/542,239

Input Set : A:\1181-8 PCT US.SeqList.txt
Output Set: N:\CRF4\07252005\J542239.raw

Output Set: N:\CRF4\07252005\J542239.raw 69 <211> LENGTH: 17 70 <212> TYPE: PRT 71 <213> ORGANISM: Artificial sequence 73 <220> FEATURE: 74 <223> OTHER INFORMATION: peptide 75 <400> SEQUENCE: A	
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DATE: 07/25/2005 TIME: 11:54:35

PATENT APPLICATION: US/10/542,239

Input Set : A:\1181-8 PCT US.SeqList.txt
Output Set: N:\CRF4\07252005\J542239.raw

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/542,239

DATE: 07/25/2005 TIME: 11:54:36

Input Set : A:\1181-8 PCT US.SeqList.txt
Output Set: N:\CRF4\07252005\J542239.raw

L:13 M:270 C: Current Application Number differs, Replaced Application Number L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date